



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/852,910A
Source: 1600
Date Processed by STIC: 10/20/2003

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OCT 21 2003
TECH CENTER 1600/2900

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.1 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/2003):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crÿstal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/2003



1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/852,910A

DATE: 10/20/2003

TIME: 07:49:21

Input Set : A:\2661-101.txt

Output Set: N:\CRF4\10202003\I852910A.raw

3 <110> APPLICANT: GILCHRIST, ANNENETTE
 4 HAMM, HEIDI
 6 <120> TITLE OF INVENTION: METHOD FOR IDENTIFYING MODULATORS OF G PROTEIN COUPLED
 RECEPTOR

7 SIGNALING
 9 <130> FILE REFERENCE: 2661-101
 11 <140> CURRENT APPLICATION NUMBER: US 09/852910A
 12 <141> CURRENT FILING DATE: 2001-05-11
 14 <150> PRIOR APPLICATION NUMBER: US 60/275472
 15 <151> PRIOR FILING DATE: 2001-03-14
 17 <160> NUMBER OF SEQ ID NOS: 271
 19 <170> SOFTWARE: PatentIn version 3.2
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 4
 23 <212> TYPE: PRT
 24 <213> ORGANISM: Mammal
 28 <220> FEATURE:
 29 <221> NAME/KEY: misc_feature
 30 <222> LOCATION: (1)..(4)
 31 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
 33 <220> FEATURE:
 34 <221> NAME/KEY: misc_feature
 35 <222> LOCATION: (1)..(4)
 36 <223> OTHER INFORMATION: MBP-G11
 38 <400> SEQUENCE: 1

W--> 40 Xaa Xaa Xaa Xaa

41 1
 44 <210> SEQ ID NO: 2
 45 <211> LENGTH: 11
 46 <212> TYPE: PRT
 47 <213> ORGANISM: Homo sapiens
 49 <400> SEQUENCE: 2
 51 Leu Gln Leu Asn Leu Lys Glu Tyr Asn Leu Val
 52 1 5 10
 55 <210> SEQ ID NO: 3
 56 <211> LENGTH: 4
 57 <212> TYPE: PRT
 58 <213> ORGANISM: Mammal
 60 <220> FEATURE:
 61 <221> NAME/KEY: misc_feature
 62 <222> LOCATION: (1)..(4)
 63 <223> OTHER INFORMATION: PAR-13
 66 <400> SEQUENCE: 3
 68 Val Arg Pro Ser

Does Not Comply
 Corrected Diskette Needed

See pp. 6-7

RAW SEQUENCE LISTING

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69 1
72 <210> SEQ ID NO: 4
73 <211> LENGTH: 11
74 <212> TYPE: PRT
75 <213> ORGANISM: Artificial Sequence
77 <220> FEATURE:
78 <223> OTHER INFORMATION: Gq peptide library sequence
81 <400> SEQUENCE: 4
83 Leu Gln Leu Asn Arg Asn Glu Tyr Tyr Leu Val
84 1 5 10
87 <210> SEQ ID NO: 5
88 <211> LENGTH: 4
89 <212> TYPE: PRT
90 <213> ORGANISM: Mammal
92 <220> FEATURE:
93 <221> NAME/KEY: misc_feature
94 <222> LOCATION: (1)..(4)
95 <223> OTHER INFORMATION: PAR-23
98 <400> SEQUENCE: 5
100 Leu Ser Arg Ser
101 1
104 <210> SEQ ID NO: 6
105 <211> LENGTH: 11
106 <212> TYPE: PRT
107 <213> ORGANISM: Artificial Sequence
109 <220> FEATURE:
110 <223> OTHER INFORMATION: Gq peptide library sequence
113 <400> SEQUENCE: 6
115 Leu Gln Gln Lys Leu Lys Glu Tyr Ser Leu Val
116 1 5 10
119 <210> SEQ ID NO: 7
120 <211> LENGTH: 4
121 <212> TYPE: PRT
122 <213> ORGANISM: Mammal
124 <220> FEATURE:
125 <221> NAME/KEY: misc_feature
126 <222> LOCATION: (1)..(4)
127 <223> OTHER INFORMATION: PAR-33
130 <400> SEQUENCE: 7
132 Leu Ser Thr Asn
133 1
136 <210> SEQ ID NO: 8
137 <211> LENGTH: 11
138 <212> TYPE: PRT
139 <213> ORGANISM: Artificial Sequence
141 <220> FEATURE:
142 <223> OTHER INFORMATION: Gq peptide library sequence
145 <400> SEQUENCE: 8
147 Leu His Leu Asn Leu Lys Glu Tyr Asn Leu Val

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Input Set: A:\2661-101.txt
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148 1 5 10
151 <210> SEQ ID NO: 9
152 <211> LENGTH: 4
153 <212> TYPE: PRT
154 <213> ORGANISM: Mammal
156 <220> FEATURE:
157 <221> NAME/KEY: misc_feature
158 <222> LOCATION: (1)..(4)
159 <223> OTHER INFORMATION: PAR-34
162 <400> SEQUENCE: 9
164 Leu Pro Gln Met
165 1
168 <210> SEQ ID NO: 10
169 <211> LENGTH: 11
170 <212> TYPE: PRT
171 <213> ORGANISM: Artificial Sequence
173 <220> FEATURE:
174 <223> OTHER INFORMATION: Gq peptide library sequence
177 <400> SEQUENCE: 10
179 Gln Arg Leu Asn Val Gly Glu Tyr Asn Leu Val
180 1 5 10
183 <210> SEQ ID NO: 11
184 <211> LENGTH: 4
185 <212> TYPE: PRT
186 <213> ORGANISM: Mammal
188 <220> FEATURE:
189 <221> NAME/KEY: misc_feature
190 <222> LOCATION: (1)..(4)
191 <223> OTHER INFORMATION: PAR-45
194 <400> SEQUENCE: 11
196 Ser Arg His Thr
197 1
200 <210> SEQ ID NO: 12
201 <211> LENGTH: 11
202 <212> TYPE: PRT
203 <213> ORGANISM: Artificial Sequence
205 <220> FEATURE:
206 <223> OTHER INFORMATION: Gq peptide library sequence
209 <400> SEQUENCE: 12
211 Leu Arg Leu Asn Gly Lys Glu Leu Asn Leu Val
212 1 5 10
215 <210> SEQ ID NO: 13
216 <211> LENGTH: 11
217 <212> TYPE: PRT
218 <213> ORGANISM: Homo sapiens
220 <400> SEQUENCE: 13
222 Gln Arg Met His Leu Arg Gln Tyr Glu Leu Leu
223 1 5 10
226 <210> SEQ ID NO: 14

RAW SEQUENCE LISTING

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Input Set : A:\2661-101.txt
 Output Set: N:\CRF4\10202003\I852910A.raw

227 <211> LENGTH: 67
 228 <212> TYPE: DNA
 229 <213> ORGANISM: Artificial Sequence
 231 <220> FEATURE:
 232 <223> OTHER INFORMATION: G alpha t library construct
 235 <220> FEATURE:
 236 <221> NAME/KEY: misc_feature
 237 <222> LOCATION: (9)..(10) /
 238 <223> OTHER INFORMATION: n is a, c, g, or t
 240 <220> FEATURE:
 241 <221> NAME/KEY: misc_feature /
 242 <222> LOCATION: (12)..(13) /
 243 <223> OTHER INFORMATION: n is a, c, g, or t
 245 <220> FEATURE:
 246 <221> NAME/KEY: misc_feature /
 247 <222> LOCATION: (15)..(16) /
 248 <223> OTHER INFORMATION: n is a, c, g, or t
 250 <220> FEATURE:
 251 <221> NAME/KEY: misc_feature
 252 <222> LOCATION: (18)..(19) /
 253 <223> OTHER INFORMATION: n is a, c, g, or t
 255 <400> SEQUENCE: 14
 W--> 256 gaggtggtnn knnnnnnnn attcgtgaaa acttaaaaga ttgtggtcgt ttctaaactaa 60
 258 gtaaaaggc 67
 261 <210> SEQ ID NO: 15
 262 <211> LENGTH: 11
 263 <212> TYPE: PRT
 264 <213> ORGANISM: Homo sapiens
 266 <400> SEQUENCE: 15
 268 Ile Lys Glu Asn Leu Lys Asp Cys Gly Leu Phe
 269 1 5 10
 272 <210> SEQ ID NO: 16
 273 <211> LENGTH: 33
 274 <212> TYPE: DNA
 275 <213> ORGANISM: Homo sapiens
 277 <400> SEQUENCE: 16
 278 atcaaggaga acctgaaaga ctgcggcctc ttc 33
 281 <210> SEQ ID NO: 17
 282 <211> LENGTH: 11
 283 <212> TYPE: PRT
 284 <213> ORGANISM: Homo sapiens
 286 <400> SEQUENCE: 17
 288 Ile Lys Asn Asn Leu Lys Asp Cys Gly Leu Phe
 289 1 5 10
 292 <210> SEQ ID NO: 18
 293 <211> LENGTH: 33
 294 <212> TYPE: DNA
 295 <213> ORGANISM: Homo sapiens
 297 <400> SEQUENCE: 18

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/852,910A

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Input Set : A:\2661-101.txt
Output Set: N:\CRF4\10202003\I852910A.raw

298 ataaaaaaaata atctaaaaaga ttgtggtctc ttc 33
 301 <210> SEQ ID NO: 19
 302 <211> LENGTH: 11
 303 <212> TYPE: PRT
 304 <213> ORGANISM: Artificial Sequence
 306 <220> FEATURE:
 307 <223> OTHER INFORMATION: G alpha i 1/2 sequence in random order
 310 <400> SEQUENCE: 19
 312 Asn Gly Ile Lys Cys Leu Phe Asn Asp Lys Leu
 313 1 5 10
 316 <210> SEQ ID NO: 20
 317 <211> LENGTH: 33
 318 <212> TYPE: DNA
 319 <213> ORGANISM: Artificial Sequence
 321 <220> FEATURE:
 322 <223> OTHER INFORMATION: G alpha i 1/2 sequence in random order
 325 <400> SEQUENCE: 20
 326 aacggcatca agtgcctttt caacgacaag ctg 33
 329 <210> SEQ ID NO: 21
 330 <211> LENGTH: 11
 331 <212> TYPE: PRT
 332 <213> ORGANISM: Homo sapiens
 334 <400> SEQUENCE: 21
 336 Ile Lys Asn Asn Leu Lys Glu Cys Gly Leu Tyr
 337 1 5 10
 340 <210> SEQ ID NO: 22
 341 <211> LENGTH: 33
 342 <212> TYPE: DNA
 343 <213> ORGANISM: Homo sapiens
 345 <400> SEQUENCE: 22
 346 ataaaaaaca acttaaagga atgtggactt tat 33
 349 <210> SEQ ID NO: 23
 350 <211> LENGTH: 11
 351 <212> TYPE: PRT
 352 <213> ORGANISM: Homo sapiens
 354 <400> SEQUENCE: 23
 356 Ile Ala Lys Asn Leu Arg Gly Cys Gly Leu Tyr
 357 1 5 10
 360 <210> SEQ ID NO: 24
 361 <211> LENGTH: 33
 362 <212> TYPE: DNA
 363 <213> ORGANISM: Homo sapiens
 365 <400> SEQUENCE: 24
 366 atcgccaaaa acctgcgggg ctgtggactc tac 33
 369 <210> SEQ ID NO: 25
 370 <211> LENGTH: 11
 371 <212> TYPE: PRT
 372 <213> ORGANISM: Homo sapiens
 374 <400> SEQUENCE: 25

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<210> SEQ ID NO 43
<211> LENGTH: 20
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: : see p. 7 for error explanation
<400> SEQUENCE: 43

gatccggccgc caccatggga

20

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/852,910A

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TIME: 07:49:22

Input Set : A:\2661-101.txt
Output Set: N:\CRF4\10202003\I852910A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,2,3,4
Seq#:14; N Pos. 9,10,12,13,15,16,18,19
Seq#:118; N Pos. 9,10,12,13,15,16,18,19
Seq#:119; N Pos. 9,10,12,13,15,16,18,19
Seq#:120; N Pos. 9,10,12,13,15,16,18,19
Seq#:121; N Pos. 9,10,12,13,15,16,18,19
Seq#:122; N Pos. 9,10,12,13,15,16,18,19
Seq#:123; N Pos. 9,10,12,13,15,16,18,19

Use of <220> Feature(NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence"
or "Unknown". Please explain source of genetic material in <220> to <223>
section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32)
(Sec.1.823 of new Rules)

Seq#:43

no explanation

VERIFICATION SUMMARYPATENT APPLICATION: **US/09/852,910A**

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Input Set : **A:\2661-101.txt**Output Set: **N:\CRF4\10202003\I852910A.raw**

L:40 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:256 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:553 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:43, <213>
ORGANISM:Artificial Sequence
L:553 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:43, <213>
ORGANISM:Artificial Sequence
L:553 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:43, Line#:553
L:1530 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:118 after pos.:0
L:1565 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:119 after pos.:0
L:1600 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:120 after pos.:0
L:1635 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:121 after pos.:0
L:1670 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:122 after pos.:0
L:1705 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:123 after pos.:0